

**AMENDMENTS TO THE SPECIFICATION**

Please amend the section of paragraphs beginning on page 5, line 6, and ending on page 5, line 14, as follows:

~~Fig. 1 illustrates~~ Figs. 1 and 2 illustrate the homology of base sequences of the IGR-IRES region of CrP-like viruses. PSIV is SEQ ID NO:13; HiPV is SEQ ID NO:14; DCV is SEQ ID NO:15; CrPV is SEQ ID NO:16; TrV is SEQ ID NO:17; BQCV is SEQ ID NO:18; RhPV is SEQ ID NO: 19.

Fig. 2 is a continuation of Fig. 1. The “•” symbol in the drawing indicates the end of the IGR-IRES region and the beginning of a virus coat protein encoding region for each sequence.

Fig. 3 illustrates a secondary structure predicted by computer program MFOLD.

Fig. 4 shows that the higher-order structure of the IGR-IRES region of CrP-like viruses is conserved. Small bold letters indicate stem structure-forming bases, and large bold letters indicate bases involved in the base pair formation of the pseudoknot. PSIV is SEQ ID NO:13; HiPV is SEQ ID NO:14; DCV is SEQ ID NO:15; CrPV is SEQ ID NO:16; TrV is SEQ ID NO:17; BQCV is SEQ ID NO:18; RhPV is SEQ ID NO: 19. For each sequence, the IGR-IRES region ends and the virus coat protein translation region begins where indicated on the drawing.

U.S. Patent Application No. 10/088,750  
Amendment dated October 7, 2005  
Reply to Office Action of July 8, 2005

**AMENDMENT TO THE SEQUENCE LISTING**

Please delete the entire sequence listing that is currently in the application and insert the substitute sequence listing that is attached hereto as a separate sheet.